

## AMENDMENTS TO THE SPECIFICATION

Amend the paragraph beginning at page 24, line 16, as follows.

Figure 3C is an alignment of *eri-1* and other members of the DEDDh family of nucleases (~~SEQ ID NOS:12, 13, 14, 15, 16, 17, 18, and 19~~) (SEQ ID NOS:12-35). This alignment was generated by the Clustal program. Shown are three conserved motifs (I, II, and III) found within the approximately 200 amino acid nuclease domain of the DEDDh superfamily of nucleases. Asterisks indicate highly conserved active site residues. Dark shading within motifs I and III indicates highly conserved residues within the DEDDh superfamily of nucleases. Lighter shading within motif II indicates substituted residues within the *eri-1* DEDDh subfamily of nucleases. A randomly chosen DEDDh nuclease domain (AAH051864) was included for reference purposes.

Insert the Sequence Listing submitted with the concurrently filed Statement under 37 C.F.R. § 1.821 at the end of the specification.